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1 [Bioinformatics—an introduction for computer scientists](#)

[Jacques Cohen](#)

June 2004 **Computing Surveys (CSUR)**, Volume 36 Issue 2

Publisher: ACM

Full text available: [Pdf](#) (261.56 KB)

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Bibliometrics: Downloads (6 Weeks): 185, Downloads (12 Months): 1183, Citations: 1

The article aims to introduce computer scientists to the new field of bioinformatics from the needs of biologists to utilize and help interpret the vast amount of data being gathered in genomic research---and ...

Keywords: DNA, Molecular cell biology, RNA and protein structure, alignment, modeling, computer, dynamic programming, hidden-Markov-models, motif discovery, sequences, phylogenetic trees

2 [A history of Haskell: being lazy with class](#)

[Paul Hudak](#), [John Hughes](#), [Simon Peyton Jones](#), [Philip Wadler](#)

June 2007 **HOPL III**: Proceedings of the third ACM SIGPLAN conference on Haskell languages

Publisher: ACM

Full text available: [Mov](#) (56:47 MIN), [Pdf](#) (1.15 MB) Additional Information: [full citation](#), [abstract](#), [bibliometrics](#)

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This paper describes the history of Haskell, including its genesis and prior implementations and tools, and applications and impact.

3 [Communications of the ACM: Volume 51 Issue 7](#)

July 2008 **Communications of the ACM**

Publisher: ACM

Full text available: [Digital Edition](#), [Pdf](#) (6.54 MB) Additional Information: [full citation](#), [abstract](#), [bibliometrics](#)

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4 [Communications of the ACM: Volume 51 Issue 12](#)

December 2008 **Communications of the ACM**

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